



# MARK YOUR CALENDARS: DECEMBER 13-14, 2012 CONFERENCE

## *Toward the 3D Virtual Cell*

UNIVERSITY OF CALIFORNIA, SAN DIEGO

Translational biology is faced with cyber-infrastructure challenges related to data accessibility, software development, software reuse and sustainability on scales not seen before. We are challenged to creating new algorithm development and analysis tools that can operate on an increasingly large, diverse, complex and widely distributed body of digital biological data.

### WE ARE EXCITED TO HOST THE FOLLOWING SPEAKERS

Theodore Alexandrov, <i>Universität Bremen</i>	Erich Huang, <i>Duke Institute for Genome Sciences &amp; Policy</i>	Lucila Ohno-Machado, <i>UC San Diego</i>
Chandrajit Bajaj, <i>University of Texas</i>	Peter Hunter, <i>University of Auckland</i>	Bernhard Palsson, <i>UC San Diego</i>
Chaitan Baru, <i>UC San Diego</i>	Graham Johnson, <i>UC San Francisco</i>	Steven Peltier, <i>UC San Diego</i>
Donald Blumenthal, <i>University of Utah</i>	Peter Karp, <i>SRI International</i>	Ramesh Rao, <i>UC San Diego</i>
Blake Borgeson, <i>University of Texas, Austin</i>	Thomas Knudsen, <i>EPA</i>	Bing Ren, <i>UC San Diego</i>
Phil Bourne, <i>UC San Diego</i>	Eugene Kolker, <i>Seattle Children's Research Center</i>	Daniel Russel, <i>UC San Francisco &amp; Google</i>
Wah Chiu, <i>Baylor College of Medicine</i>	Leslie Loew, <i>University of Connecticut Health Center</i>	Hebert Sauro, <i>University of Washington</i>
Tim Clark, <i>Harvard University</i>	Maryann Martone, <i>UC San Diego</i>	Terry Sejnowski, <i>UC San Diego</i>
Phillip Compeau, <i>UC San Diego</i>	Andrew McCulloch, <i>UC San Diego</i>	Nigam Shah, <i>Stanford University</i>
Markus Covert, <i>Stanford University</i>	Martin Meier-Schellersheim, <i>NIH</i>	James Sluka, <i>Indiana University</i>
Anita de Waard, <i>Elsevier</i>	Marcel Oberlaender, <i>Max Planck Institute for Biological Cybernetics</i>	Maciej Swat, <i>Indiana University</i>
Pieter Dorrestein, <i>UC San Diego</i>	Arthur Olson, <i>The Scripps Research Institute</i>	Susan Taylor, <i>UC San Diego</i>
Mark Ellisman, Ph.D., <i>UC San Diego</i>		Wei Wang, , <i>UC San Diego</i>
Yolanda Gil, <i>USC</i>		Nancy Wilkins-Diehr, <i>UC San Diego</i>
James Glazier, <i>Indiana University</i>		Ross Whitaker, <i>University of Utah</i>
David Goodsell, <i>The Scripps Research Institute</i>		

### AREAS TO BE EXPLORED

- Structure-based modeling
- Visualization
- Whole cell modeling
- Pathway modeling
- Neurological modeling
- Ontologies and Tools for modeling
- Modeling big data
- Software engineering workflows for modeling
- New modes of education and dissemination

### CONFERENCE OUTCOMES

- Recognizing the scientific challenges
- Identifying today's tools for cellular modeling
- Defining needed developments in algorithms, visualization, software paradigms, workflow development, education and more...

### STEERING COMMITTEE

- Philip Bourne, Ph.D. , PI, *UC San Diego*
- Eugene Kolker, Ph.D. Co-PI, *Seattle Children's Research Center*
- Peter Arzberger, Ph.D. , *UC San Diego*
- Helen Berman, Ph.D. , *Rutgers University*
- Donald Blumenthal, Ph.D., *University of Utah*
- Mark Ellisman, Ph.D., *UC San Diego*
- Andrew McCullough, Ph.D., *UC San Diego*
- Bernhard Palsson, Ph.D., *UC San Diego*
- Ramesh Rao, Ph.D., *UC San Diego*
- Susan Taylor, Ph.D. , *UC San Diego*

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